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Listing first 45 summaries
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Copyright (c) 1993 - 2000 Compugen Ltd.
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CH62_SYNVU
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YCX7_CHLRE
IM08_ARATH
LPPB_HUMAN
SECR_PIG
YQ1B_ECOLI
HBAT_HUMAN
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PHCB_SYNPW
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VMS1_MOUSE
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ANNEXIN 1.
ANNEXIN 2.
INTERCHAIN CROSS-LINK (BY SIMILARITY).
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01-NOV-1995 (Rel. 32, Last sequence update)
01-NOV-1995 (Rel. 32, Last annotation update)
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P20384;
                                                                                                                                                                                         PROSITE;
                                                                                                                                                                                                                                                                                                                            the European Bioinformatics Institute. There are no restrictions use by non-profit institutions as long as its content is in modified and this statement is not removed. Usage by and for contents
                                                                                                                                                                                                                                                                                                                                                                                                                      Submitted (SEP-1993) to the EMBL/GenBank/DDBJ databases.
-!- FUNCTION: BIN3 IS A POTENTIAL DNA INVERTASE.
-!- SIMILARITY: BELONGS TO THE "RESOLVASE" FAMILY OF SITE-SPECIFIC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              STRAIN-NCTC 9789;
MEDLINE-90060039; PubMed-2555186;
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Bacteria; Firmicutes;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Potential DNA-invertase BIN3 (Transposon TN552).
                                                                                                                                                                                                                                                                                                or send an email to license@isb-sib.ch).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                         STRAIN-NCTC 9789;
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ACT_SITE
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pfam; pF02796; HTH_7; 1.
pfam; pF00239; resolvase; 1.
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118 AMVSE 122
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                         1 AMVSE 5
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DNA integration; DNA invertase.
9 9 TRANSIENT COVALENT LINKAGE TO DNA DURING
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Best Local
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MEDLINE=98013059; PubMed=9353010;
Bosse J.T., Macinnes J.I.;
"Genetic and biochemical analyses of Actinobacillus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        URED_ACTPL STANDARD; PRT; 270 AA. 054425; 05-DEC-1998 (Rel. 37, Created) 15-DEC-1998 (Rel. 37, Last sequence update) 30-MAY-2000 (Rel. 39, Last annotation update)
                                                                                                                                                                                                                      AN12_COLLI STANDARD; PRT; 343 AA.
Q92040; Q92041;
01-NOV-1997 (Rel. 35, Created)
01-NOV-1997 (Rel. 35, Last sequence update)
16-OCT-2001 (Rel. 40, Last annotation update)
Annexin I, isoform p37 (Lipocortin I) (Calpactin II) (Chromobindin 9)
(Phospholipase A2 inhibitory protein).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               use by non-profit institutions as long as its content is in no way modified and this statement is not removed. Usage by and for commercial entities requires a license agreement (See http://www.isb-sib.ch/announce/
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           This SWISS-PROT entry is copyright. It is produced through a collaboration between the Swiss Institute of Bioinformatics and the EMBL outstation the European Bioinformatics Institute. There are no restrictions on its burgers and the EMBL outstation its
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Actinobacillus pleuropneumoniae (Haemophilus pleuropneumoniae). Bacteria; Proteobacteria; gamma subdivision; Pasteurellaceae;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ACTPL
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                                                                          SEQUENCE FROM N.A., AND SEQUENCE OF 13-50 STRAIN=WHITE CARNEAU; TISSUE=Cropsac; MEDLINE=92406850; PubMed=1388165;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              NCBI_TaxID=715;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Actinobacillus.
           Haigler H.T., Mangili J.A., Gao Y., Jones J., Horseman N.D.; "Identification and characterization of columbid annexin Icp Insights into the evolution of annexin I phosphorylation sit J. Biol. Chem. 267:19123-19129(1992).
                                                                                                                                                            Archosauria;
                                                                                                                                                                          Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
                                                                                                                                                                                            Columba livia (Domestic pigeon).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    SEQUENCE
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Pfam; PF01774; UreD;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     InterPro; IPR002669; UreD.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   EMBL; U89957; AAC00065.1; -.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   or send an email to license@isb-sib.ch).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Infect. Immun. 65:4389-4394(1997).
-!- FUNCTION: PROBABLY FACILITATES NICKEL INCORPORATION.
-!- SIMILARITY: BELONGS TO THE URED FAMILY.
                                                                                                                                           NCBI_TaxID=8932;
                                                                                                                                                                                                                                                                                                                                                                                                     213 AMVSE
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Pred. No.
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RESULT ANXILHE ANXILHE
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Best Local
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MOD_RES
CONFLICT
SEQUENCE
                                                                                                                                 P04083;
P04083;
P01-NOV-1986 (Rel. 03, Created)
01-NOV-1986 (Rel. 03, Last sequence update)
01-MAR-2002 (Rel. 41, Last annotation update)
Annexin I (Lipocortin I) (Calpactin II) (Chromobindin 9) (P35)
(Phospholipase A2 inhibitory protein).
ANXA1 OR ANXI OR LPC1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            EMBL;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               "Structural and functional diversenceding cp37 and cp35 genes."; Gene 143:179-186(1994).
                                                                                   Eukaryota; Metazoa;
Mammalia; Eutheria;
                                                                                                                                                                                                                                                                                                                 HUMAN
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      PRINTS;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          InterPro; IPR001464; Annexin
Pfam; PF00191; annexin; 4.
PRINTS; PR00196; ANNEXIN.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        HSSP;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           EMBL; M91008; AAA49447.1; ALT_INIT.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      or send an email to license@isb-sib.ch).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         entities requires a license agreement (See http://www.isb-sib.ch/announce/
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      MEDLINE-86146879;
                         SEQUENCE FROM N.A.
                                                                                                                                                                                                                                                                                            ANX1_HUMAN
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                                                                NCBI_TaxID=9606;
                                                                                                                          Homo sapiens (Human)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          FUNCTION: CALCIUM/PHOSPHOLIPID-BINDING PROTEIN WHICH PROMOTES MEMBRANE FUSION AND IS INVOLVED IN EXOCYTOSIS. THIS PROTEIN REGULATES PHOSPHOLIPASE AZ ACTIVITY. IT SEEMS TO BIND FROM TWE FOUR CALCIUM IONS WITH HIGH AFFINITY.

PTM: PHOSPHORYLATED BY PROTEIN KINASE C AND EPIDERMAL GROWTH
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              FACTOR RECEPTOR/KINASE.
PTM: THE N-TERMINUS IS BLOCKED.
MISCELLANEOUS: IN PIGEONS, TWO ISOFORMS OF
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               BY THE DIFFERENTIALLY REGULATED
                                                                                                                                                                                                                                                                                                                                                                                                                                       AMVSE 5
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een the Swiss Institute of Bioinformatics and the EN
                                                                                                                                                                                                                                                                                                                                                                                                 AMVSE
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203
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1BO9.
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; ANX; 4.
                                                                                                                                                                                                                                                                                            STANDARD;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            inhibitor;
      PubMed=2936963
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179
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                                                                                                       Chordata;
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ANNEXIN 4.
INTERCHAIN CROSS-LINK (BY SIMILARITY).
PHOSPHORYLATION (BY EGER).
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Pred. No.
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                                                                                      Catarrhini;
                                                                                                       Craniata; Vertebrata; Euteleostomi;
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37;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Length 343;
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autonomous folding unit.";
J. Biol. Chem. 274:2971-2977(1999).
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Kovacic R.T., Ti
"Correlation of
lipocortin I.";
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       MEDLINE=91369906; PubMed=1832554;
Kovacic R.T., Tizard R., Cate R.L., Frey A.Z., Wallner
"Correlation of gene and protein structure of rat and l
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Ramachandran K.L., Pepinsky R.B.;
"Cloning and expression of human lipocortin,
inhibitor with potential anti-inflammatory a
                                                                                                                                                                                                                                                                                                             Gao
                                                                                                                                                                                                                                                                                                                                        STRUCTURE BY NMR OF 40-112
                                                                                                                                                                                                                                                                                                                                                                  Protein
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                                                                                                                                                                                                                                                                                                                                                                                                                MEDLINE=93200922; PubMed=8453382;
                                                                                                                                                                                                                                                                                                                                                                                                                                                           Science 237:992-998(1987).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                           modifications in proteins.";
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Biemann K.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      MEDLINE=87292145; PubMed=3303336;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   "Location of sites in human lipocortin I that are phosphorylated protein tyrosine kinases and protein kinases A and C.";
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Varticovski L., Chahwala S.B., Whitman Chow E.P., Sinclair L.K., Pepinsky R.B.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             MEDLINE=88309771; PubMed=2457390;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              SEQUENCE,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Talenti A., di Rosa M., Ciliberto G.;
"Structural characterization of a biologically active
lipocortin 1 expressed in Escherichia coli.";
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Arcone R., Arpaia G., Ru
Ialenti A., di Rosa M.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               MEDLINE=93145967;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Biochemistry 30:9015-9021(1991).
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Wallner B.P., Mattaliano R.J., Sinclair L.K., Foeller C., Cho
                                                                                                                                                                                                                                                                                            "NMR solution structure of domain 1 of human annexin I shows
                                                                                                                                                                                                                                                                                                                      MEDLINE=99115644; PubMed=9915835;
                                                                                                                                                                                                                                                                                                                                                                                                                              X-RAY CRYSTALLOGRAPHY (2.5 ANGSTROMS).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          DIMERIC
                                                                                                                                                                                                                                                                                                                                                                                   "Crystal
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       "Characterization by tandem mass
                                                                                                             MEMBRANE FUSION AND IS INVOLVED IN EXOCYTOSIS. THIS PROTEIN REGULARTES PHOSPHOLIPASE A2 ACTIVITY. IT SEEMS TO BIND FROM TO FOUR CALCIUM IONS WITH HIGH AFFINITY.

SUBUNIT: IN PLACEMYA 20% OF THE PROTEIN IS FOUND AS A COVALES CROSS-LINKED DIMER WHICH IS GENERATED BY A TRANSGLUTAMINASE.

CONSTAINS FOUR HOMOLOGOUS REPEATS WITH A CONSENSUS SEQUENCE COMMON TO ALL ANNEXIN PROTEINS. A PAIR OF THESE REPIMAY FORM ONE BINDING SITE FOR CALCIUM AND PHOSPHOLIPID.

MAY FORM ONE BINDING SITE FOR CALCIUM AND PHOSPHOLIPID.

PIM: PHOSPHORYLATED BY PROTEIN KINASE C AND EPIDERMAL GROWTH
                                                       INHIBITORY ACTIVITY.
SIMILARITY: BELONGS TO THE ANNEXIN
DATABASE: NAME=Annexin I home page:
                                                                                                                                                                                                                                                                                                        J., Li Y., Yan H.;
                                                                                                                                                                                                                                                 FUNCTION: CALCIUM/PHOSPHOLIPID-BINDING PROTEIN
                                       WWW="http://www.mds.qmw.a.c.uk/biopharm/annI.html"
                                                                                                  FACTOR RECEPTOR/KINASE. PHOSPHORYLATION RESULTS
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01-MAR-1989 (Rel. 10, Last sequence update)
16-OCT-2001 (Rel. 40, Last annotation update)
Annexin I (Lipocortin I) (Calpactin II) (Chromobindin (Phospholipase A2 inhibitory protein).
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                                 MEDLINE=91301692; PubMed=1676980;
Horlick K.R., Cheng I.C., Wong W.T., Wakeland E.K.,
"Mouse lipocortin I gene structure and chromosomal a
duplication and the origins of a gene family.";
Genomics 10:365-374(1991).
                                                                                                                                  SEQUENCE FROM
                                                                                                                                                                                                                               Sakata T., Iwagami
                                                                                                                                                                                                                                                     MEDLINE-89098333;
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                                                                                                                                                                                                               Teraoka H.;
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PDB; 1BO9; 19-AUG-98.
PHCI-2DPAGE; P04083;
MIM; 151690; -.
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-i- FUNCTION: CALCIUM,PHOSPHOLIPID-BINDING PROTEIN WHICH PROMOTES MEMBRANE FUSION AND IS INVOLVED IN EXOCYTOSIS. THIS PROTEIN REGULARES PHOSPHOLIPASE A2 ACTIVITY. IT SEEMS TO BIND FROM TWO TO
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PTM.: PHOSPHORYLATED BY PROTEIN KINASE C AND EPIDERMAL GROWPTM: PROCEPTOR/KINASE. PHOSPHORYLATION RESULTS IN LOSS OF INHIBITORY ACTIVITY (BY SIMILARITY).

SIMILARITY: BELONGS TO THE ANNEXIN FAMILY.
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P07150; Q64664;
01-APR-1988 (Rel
                                                                                                                                                     EMBL;
                                                                                                                                                                                                                                                                                                                                                                                                                                          Biochemistry 30:9015-9021(1991).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     STRAIN-SPRAGUE-DAWLEY; TISSUE-Liver;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       "Molecular cloning and expression in Escherichia coli of the cDNA coding for rat lipocortin I (calpactin II).";
Gene 65:141-147(1988).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       RAT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Kovacic R.T., Tizard R., Cate R.L., Frey A.Z., Wallner B.P., "Correlation of gene and protein structure of rat and human."
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      SEQUENCE FROM N.A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Shimizu Y., Takabayashi
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           MEDLINE=91369906; PubMed=1832554;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  01-APR-1988 (Rel. 07, Last sequence update)
16-OCT-2001 (Rel. 40, Last annotation update)
Annexin I (Lipocortin I) (Calpactin II) (Chromobindin 9) (Phospholipase A2 inhibitory protein).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       MEDLINE=88284376; PubMed=2969352;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Tamaki M., Nakamura E., Nishikubo C., Sakata T., "Rat lipocortin I cDNA.";
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             MEDLINE=88015620; PubMed=2958780;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                            .lpocortin I
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Mammalia; Eutheria;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Eukaryota; Metazoa;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        SEQUENCE FROM N.A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Rattus norvegicus (Rat)
                                                                                                                                                                                                                                                                                                                  FACTOR RECEPTOR/KINASE. PHOSPHORYLATION RESULTS INHIBITORY ACTIVITY (BY SIMILARITY). SIMILARITY: BELONGS TO THE ANNEXIN FAMILY.
                                                                                                                                                                                                                                                                                                                                                     REGULATES PHOSPHOLIPASE AZ ACTIVITY, IT SEEMS TO BIND FROM TWO TO FOUR CALCIUM IONS WITH HIGH APPINITY.

DOMAIN: CONTAINS FOUR HOMOLOGOUS REPEATS WITH A CONSENSUS SEQUENCE COMMON TO ALL ANNEXIN PROTEINS. A PAIR OF THESE REPEATS PAMPETORM ONE BINDING SITE FOR CALCIUM AND PHOSPHOLIPID.

PTM: PHOSPHORYLATED BY PROTEIN KINASE C AND EPIDERMAL GROWTH
                           JT0303;
                                                                                                                                                                                                                                                                                                                                                                                                                                 MEMBRANE FUSION AND IS INVOLVED IN EXOCYTOSIS. THIS PROTEIN
                                      $57468;
$57470;
$57472;
$57474;
$57474;
                                                                                                                                                  S57447;
S57450;
                                                                                                                                                                           M19967;
S57478;
               P04083;
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IPR001464; Annexin
                        LURT 1
                                                                                                                                                 AAB19866.1;
AAB19866.1;
                                                                                                                                                                                      AAA40861.1; -.
               1809
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                                      AAB19866.1;
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Rodentia;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Created)
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Sciurognathi;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Vertebrata; Euteleostomi;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Muridae; Murinae; Rattus.
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tent is in
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                                                                                                                                                                                                                                                                            EMBL outstation
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C. -I-FUNCTION: CALCIUM/PHOSPHOLIPID-BINDING PROTEIN WHICH PKOMUTES
C. MEMBRANE FUSION AND IS INVOLVED IN EXOCYTOSIS.
C. -I-FUNCTION: AGGREGATION FACTOR (AF)-ASSOCIATED CALLBACTIN AND PLASMA-
C. MEMBRANE-BOUND PHOSPHOLIPIDS ARE INVOLVED IN CALCIUM DEPENDENT
C. TELL-CELL AGGREGATION IN SPONGES, CALPACTIN BINDS PHOSPHOLIPIDS.
C. IT BINDS TWO CALCIUM IONS/MOL WITH LOW AFFINITY IN THE ABSENCE OF
C. PHOSPHOLIPIDS, AND WITH HIGH AFFINITY IN THEIR PRESENCE.
C. SEQUENCE COMMON TO ALL ANNEXIN PROTEINS. A PAIR OF THESE REPEATS
C. SEQUENCE COMMON TO ALL ANNEXIN PROTEINS. A PAIR OF THESE REPEATS
C. SEQUENCE COMMON TO ALL ANNEXIN PROTEINS. A PAIR OF THESE REPEATS
C. PAY FORM ONE BINDING SITE FOR CALCIUM AND PHOSPHOLIPID.
C. PIM: PHOSPHORYLATED BY PROTEIN KINASE C AND EPIDERMAL GROWTH
FACTOR RECEPTOR/KINASE. PHOSPHORYLATION RESULTS IN LOSS OF THE
C. INHIBITORY ACTIVITY (BY SIMILARITY).
C. INHIBITORY ACTIVITY (BY SIMILARITY).
C. INHIBITORY BELONGS TO THE ANNEXIN FAMILY.
C. SIMILARITY: BELONGS TO THE ANNEXIN FAMILY.
C. SIMILARITY: BELONGS TO THE ANNEXIN FAMILY.
C. BUT, ON THE BASIS OF PHYLOGENETIC STUDIES (REF. 2), IT SEEMS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ANX1
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Best Local S
Matches 5
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CONFLICT
SEQUENCE
                                                                                                                                                                                                                                                                Unpublished observations (APR-1996).
                                                                                                                                                                                                                                                                                   Mueller W
                                                                                                                                                                                                                                                                                              AGREEMENT WITH RODENT CONTAMINATION.
                                                                                                                                                                                                                                                                                                                          Unpublished observations (FEB-1996).
                                                                                                                                                                                                                                                                                                                                             Duret
                                                                                                                                                                                                                                                                                                                                                        DOUBT ON BIOLOGICAL SOURCE OF PROTEIN
                                                                                                                                                                                                                                                                                                                                                                                                       Ca2(+)-dependent cell
                                                                                                                                                                                                                                                                                                                                                                                                                             Robitzki A., Schroeder H.C., Ugarkovic D., Batel R., Mueller W.E.G.;
                                                                                                                                                                                                                                                                                                                                                                                                                                                               MEDLINE=91054405;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Eukaryota; Metazoa;
Mammalia; Eutheria;
MCBI_TaxID=69158;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              01-MAR-1992 (Rel. 21, Created)
01-OCT-1996 (Rel. 34, Last sequence update)
16-OCT-2001 (Rel. 40, Last annotation update)
                                                                                                                                                                                                                                                                                                                                                                                                            "CDNA structure and expression of calpactin, a peptide involved in
                                                                                                                                                                                                                                                                                                                                                                                                                                                                               SEQUENCE FROM N.A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Rodentia sp.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ANXA1 OR ANX1.
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SMART; SM00335; P
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Pfam; PF00191; annexin; PRINTS; PR00196; ANNEXII
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INIT_MET
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Local Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Sutheria; Rodentia; unclassified rodents.
                                                                                                                                                                                                                                                                                                                                                                                     ident cell aggregation in sponges."; 271:415-420(1990).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               345 AA;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                               PubMed=2146952;
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P -> S (IN REF. 3).
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INTERCHAIN CROSS-LINK (BY SIMILARITY).
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Pred. No.
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74508E225227F224 CRC64;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              DB 1;
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Best Local
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SEQUENCE
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INIT_MET
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             PRINTS;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Pfam; PF00191; annexin;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Bos taurus (Bovine).
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Cetartiodactyla; Ruminantia; Pecora; Bovoidea;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    01-NOV-1995 (Rel. 32, Created)
01-NOV-1995 (Rel. 32, Last sequence update)
10-NOV-1995 (Rel. 32, Last sequence update)
16-OCT-2001 (Rel. 40, Last anotation update)
Annexin I (Lipocortin I) (Calpactin II) (Chromobindin 9) (P35)
(Phospholipase A2 inhibitory protein).
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           MOD_RES
                                                                                                                                                                                                                                                                                                                 "Epitope mapping of annexin I: antibodies that compete with phospholipids and calcium recognize amino acids 42-99."; Biochem. J. 289:539-542(1993).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ANXA1 OR ANX1
                                                                                                                                                                                                                                                                                                                                                                                                                           SEQUENCE FROM N.A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                       NCBI_TaxID=9913;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Bovidae; Bovinae; Bos.
   or send an email to license@isb-sib.ch).
                 the European Bioinformatics Institute. There are no restrictions on its use by non-profit institutions as long as its content is in no way modified and this statement is not removed. Usage by and for commercial entities requires a license agreement (See http://www.isb-sib.ch/announce/
                                                                                    This SWISS-PROT entry is copyright. It is produced through a collaboration between the Swiss Institute of Bioinformatics and the EMBL outstation is the second control of the swiss institute.
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hes 5; Conserv
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                                                                                                                                  MEMBRANE FUSION AND IS INVOLVED IN EXCEPTORIS. THIS PROTEIN
REGULATES PHOSPHOLIPASE A2 ACTIVITY: IT SEEMS TO BIND FROM TWO TO
REGULATES PHOSPHOLIPASE A2 ACTIVITY: BINDS MUSCLE ACTIN.
DOMAIN: CONTAINS FOUR HOMOLOGOUS REPEATS WITH A CONSENSUS
SEQUENCE COMMON TO ALL ANNEXIN PROTEINS. A PAIR OF THESE REPEATS
MAY FORM ONE BINDING SITE FOR CALCIUM AND PHOSPHOLIPID:
MAY FORM ONE BINDING SITE FOR CALCIUM AND PHOSPHOLIPID:
MAY FORM ONE BINDING SITE FOR CALCIUM AND PHOSPHORYLATED BY PROTEIN KINASE C AND EPIDERMAL GROWTH
FACTOR RECEPTOR/KINASE, PHOSPHORYLATION RESULTS IN LOSS OF THE
INHIBITORY ACTIVITY (BY SIMILARITY).

SIMILARITY: BELONGS TO THE ANNEXIN FAMILY.
                                                                                                                                                                                                                                                                                                     FUNCTION: CALCIUM/PHOSPHOLIPID-BINDING PROTEIN WHICH PROMOTES
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ANNEXIN 3.
ANNEXIN 4.
INTERCHAIN CROSS-LINK (BY SIMILARITY).
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PHOSPHORYLATION (BY PKC) (BY SIMILARITY).
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Pred. No.
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SEQUENCE
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P19619; Q29547; Created)
01-FEB-1991 (Rel. 17, Created)
01-MAR-2002 (Rel. 41, Last sequence update)
01-MAR-2002 (Rel. 41, Last annotation update)
01-MAR-2002 (Rel. 41, Last annotation update)
Annexin I (Lipocortin I) (Calpactin II) (Chromobindin 9) (P35)
Annexin I (Lipocortin I) (Calpactin II) (Chromobindin 9) (P35)
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 membrane aggregation.";
J. MO1. Biol. 306:489-498(2001).
1- FUNCTION: CALCIUM/PHOSPHOLIPID-BINDING PROTEIN WHICH PROMOTES
-1- FUNCTION: CALCIUM/PHOSPHOLIPID IN EXOCYTOSIS. THIS PROTEIN REGULATES PHOSPHOLIPASE A2 ACTIVITY. IT SEEMS TO BIND FROM TWO TO FOUR CALCIUM IONS WITH HIGH AFFINITY.

FOUR CALCIUM IONS WITH HIGH AFFINITY BY A CONSENSUS
-1- DOMAIN: CONTAINS FOUR HOMOLOGOUS REPEATS WITH A CONSENSUS SEQUENCE COMMON TO ALL ANNEXIN PROTEINS. A PAIR OF THESE REPEATS MAY FORM ONE BINDING SITE FOR CALCIUM AND PHOSPHOLIPID.
                                                                                                                                                                                                                       De B.K., Misono K.S., Lukas T.J., Mroczkowski B., Cohen S.;
"A calcium-dependent 35-kilodalton substrate for epidermal growth
factor receptor/Kinase isolated from normal tissue.";
                                                                                                                                                                                                                                                                                                            Scemann_J., Weber K., Osborn M., Parton R.G., Gerke V. "The association of annexin I with early endosomes is Ca2+ and requires an intact N-terminal domain."; mol. Biol. Cell 7:1359-1374(1996).
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Sus scrofa
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                                                                                                                                                                                                                                                                     MEDLINE=87008618; PubMed=3020049;
                                                                                                                                                                                                                                                                                    SEQUENCE OF 13-42.
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                                                                                                                                                                                                                                                                                                                                                                                                                                 NCBI_TaxID=9823;
                                                                                                                                                                                                                                                                                                                                                                                                                                                 Mammalia; Eutheria; Cetartiodactyla;
                                                                                                                                 Rosengarth A., Gerke V., Luecke H., ^{\rm "RS} structure of full-length annexin 1 and implications
                                                                                                                                                             MEDLINE=21103721; PubMed=11178908;
                                                                                                                                                                               X-RAY CRYSTALLOGRAPHY (1.8 ANGSTROMS).
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PHOSPHORYLATION (BY PKC) (BY SIMILARITY).

CD30701636DB23A7 CRC64;
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Pred. No.
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                                                        TISSUE-Lung;
Trao F.H.C., Wen C., Hu J.;
Trace F. FUNCTION: CALCIUM/PHOSPHOLIPID-BINNING PROTEIN WHICH PROMOTES
MEMBRANE PUSION AND IS INVOLVED IN EXOCUTOSIS. THIS PROTEIN
REGULATES PHOSPHOLIPASE A2 ACTIVITY. IT SEEMS TO BIND FROM TWO TO
FOUR CALCIUM IONS WITH HIGH AFFINITY.
TO DMALN: CONTAINS FOUR HOMOLOGOUS REPEATS WITH A CONSENSUS
SEQUENCE COMMON TO ALL ANNEXIN PROTEINS. A PAIR OF THESE REPEATS
MAY FORM ONE BINDING SITE FOR CALCIUM AND PHOSPHOLIPID.
THE PTM: PHOSPHORYLATED BY PROTEIN KINASE C AND EPIDERMAL GROWTH
FACTOR RECEPTOR/KINASE. PHOSPHORYLATION RESULTS IN LOSS OF THE
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ANX1_RABIT STANDARD; PRT; 346 AA. P51662; P51662; Ol-OCT-1996 (Rel. 34, Created) Ol-OCT-1996 (Rel. 34, Last sequence update) 16-OCT-2001 (Rel. 40, Last annotation update) Annexin I (Lipocortin I) (Calpactin II) (Chromobindin 9) (P35) (Phospholipase A2 inhibitory protein).
                      <del>'</del>
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Mammalia; Eutheria;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Oryctolagus cuniculus (Rabbit).
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
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SIMILARITY: BELONGS TO THE ANNEXIN FAMILY.
                      SIMILARITY:
                   INHIBITORY ACTIVITY (BY SIMILARITY) SIMILARITY: BELONGS TO THE ANNEXIN
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         INTERCHAIN CROSS-LINK (BY SIMILARITY).
PHOSPHORYLATION (BY EGFR).
PHOSPHORYLATION (BY PKC) (BY SIMILARIT
T -> S (IN REF. 2).
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                   ANNEXIN FAMILY
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     925845C5DEBCBAB2 CRC64;
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38;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                    RESULT
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01-JUL-1993
15-JUL-1999
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       MOD_RES
SEQUENCE
                                 EMBL;
                                                                                                                                                                                                                           SEQUENCE FROM N.A.
STRAIN-H16 / ATCC 17699;
MEDLINE-93356597; PubMed-8352644;
                                                                                                                                                                             Dernedde J., Eitinger M., Friedrich B.;
"Analysis of a pleiotropic gene region involved in formation of catalytically active hydrogenases in Alcaligenes eutrophus H16."; arch. Microbiol. 159:545-553(1993).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                use by non-profit institutions as long as its content modified and this statement is not removed. Usage by an entities requires a license agreement (See http://www.isb-
                     PIR; S29978; S29978
                                                             or send an
                                                                        entities
                                                                                use by non-profit institutions as long as its content modified and this statement is not removed. Usage by an
                                                                                                            the European Bioinformatics Institute.
                                                                                                                         This SWISS-PROT entry is copyright. It is produced through a collaboration between the Swiss Institute of Bioinformatics and the EMBL outstation -
                                                                                                                                                      Arch, Microbiol, 159:545-553(1993).
                                                                                                                                                                                                                                                                                               Ralstonia
                                                                                                                                                                                                                                                                                                           Bacteria;
                                                                                                                                                                                                                                                                                                                     Alcaligenes eutrophus (Ralstonia eutropha). Plasmid megaplasmid pHG1.
                                                                                                                                                                                                                                                                                                                                                HYPD
                                                                                                                                                                                                                                                                                                                                                          Hydrogenase
                                                                                                                                                                                                                                                                                                                                                                                                               P31903;
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MOD_RES
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        PROSITE; PS00223; ANNEXIN; 3. Annexin; Calcium/phospholipid-binding; Repeat; Phosphorylation; Phospholipase A2 inhibitor.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             EMBL;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     or send an email to license@isb-sib.ch).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    This SWISS-PROT entry is copyright. It is produced through a collaboration between the Swiss Institute of Bioinformatics and the EMBL outstation - the European Bioinformatics Institute. There are no restrictions on its
            InterPro;
                                                                                                                                                                                                                                                                                NCBI_TaxID=510;
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                                    X70183; CAA49734.1;
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                                                                        requires a
          IPR002780;
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                                                             email to license@isb-sib.ch)
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346
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ilarity 100.0%;
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(Rel. 26, Last sequence update)
(Rel. 38, Last annotation updat
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38735
                                                                        license agreement
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ANNEXIN 2.
ANNEXIN 3.
ANNEXIN 4.
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Pred. No. 38;
D; Mismatches
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                                                                        (See http://www.isb-sib.ch/announce/
                                                                                                                                                                                                                                                                                                                                                                        update)
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38;
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NH10_CAEEL
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SEQUENCE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Fulton L., Waterston R.;
Submitted (JUN-1994) to the EMBL/GenBank/DDBJ databases.
-i- FUNCTION: ORPHAN NUCLEAR RECEPTOR.
-i- SUBCELLULAR LOCATION: Nuclear (Potential).
-i- SUMILARITY: BELONGS TO THE NUCLEAR HORMONE RECEPTORS FAMILY.
                                                                                                                                                                                                                                                                                                                                                                                                                                              WormPep; B0280.8; CE00815.
InterPro; IPR000536; Hormone_rec_lig.
InterPro; IPR001628; zf-C4.
Pfam; PF00104; hormone_rec; 1.
Pfam; PF00105; zf-C4; 1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                This SWISS-PROT entry is copyright. It is produced through a collaboration between the Swiss Institute of Bioinformatics and the EMBI outstation the European Bioinformatics Institute. There are no restrictions on its use by non-profit institutions as long as its content is in no way modified and this statement is not removed. Usage by and for commercial
                                                                                                                                                                                                                                                                                                                                                SMART; SM00430; HOLI; 1.
SMART; SM00399; ZnF_C4; 1.
PROSITE; PS00031; NUCLEAR_RECEPTOR;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    EMBL; U10438; HSSP; P19793;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             modified and this statement is not removed. Usage by and for commercial entities requires a license agreement (See http://www.isb-sib.ch/announce/
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              STRAIN-BRISTOL N2;
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Rhabditidae; Peloder
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Caenorhabditis elegans.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       01-NOV-1995 (Rel. 32, Created)
01-NOV-1995 (Rel. 32, Last sequence update)
30-MAY-2000 (Rel. 39, Last annotation updat
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      P41999;
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                                                                                                                                                                                                                                                                                                    Zinc-finger.
                                                                                                                                                                                                                                                                                                                             Receptor; Transcription
                                                                                                                                                                                                                                                                                                                                                                                                                        PRINTS; PR00047; STROIDFINGER
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         or send an email to license@isb-sib.ch).
                                                                                                                                                                                                                                                                                 DNA_BIND
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119
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379 AA;
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                                                                                      100.0%; ilarity 100.0%; Conservative 0
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41907 MW;
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38
78
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Pred.
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Pred. No.
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Interpro; IPR000980; SH2.
Interpro; IPR001452; SH3.
Interpro; IPR001245; Tyr_pkinase.
Pfam; PF00069; pkinase; 1.
Pfam; PF00017; SH2; I.
Pfam; PF00018; SH3; I.
Pfam; PF00018; SH3; I.
PRINTS; PR00109; TYRKINASE.
SMART; SM00250; SH3; I.
SMART; SM00250; SH3; I.
SMART; SM00219; TYFKC; I.
SMART; SM00219; TYFKC; I.
SMART; SM00219; TYFKC; I.
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SRC_RSVPA
P31693;
01-JUL-1993
DOMAIN
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BINDING
ACT_SITE
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SEQUENCE
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PROSITE;
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SEQUENCE FROM N.A.
MEDLINE=92351554; PubMed=1322589;
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DOMAIN
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            "Small deletion in v-src SH3 domain of a transformation defective mutant of Rous sarcoma virus restores wild type transforming properties.";
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01-JUL-1993 (Rel. 26, Last sequence update)
01-MAR-2002 (Rel. 41, Last annotation updat
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    HSSP; P00524; 1SPS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               EMBL; M84475; AAA42581.1; ALT_SEQ.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            or send an email to license@isb-sib.ch).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  This
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Virology 189:556-567(1992)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Rous sarcoma virus (strain PA101T).
Viruses; Retroid viruses; Retroviridae;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Tyrosine-protein kinase transforming
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                                                                                                                                                                                                        Tyrosine-protein kinase; ATP-binding; Myristate;
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SIMILARITY: CONTAINS 1 SH2 DOMAIN.
SIMILARITY: CONTAINS 1 SH3 DOMAIN.
SIMILARITY: TO OTHER PROTEIN-TYROSINE
DOMAIN. BELONGS TO THE SRC SUBFAMILY.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      FUNCTION: THIS PHOSPHOPROTEIN, REQUIRED FOR BOTH THE INITIATION AND THE MAINTENANCE OF NEOPLASTIC TRANSFORMATION, IS A PROTEIN KINASE THAT CATALYZES THE PHOSPHORYLATION OF TYROSINE RESIDUES
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  European Bioinformatics Institute.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         CATALYTIC ACTIVITY: ATP + a
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            A42994; TVFVMT.
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                                                                                                                                                                                                                                                                    PS00119; PROTEIN_KINASE_ATP;
PS00119; PROTEIN_KINASE_TYR;
PS50011; PROTEIN_KINASE_DOM;
PS50001; SH2; 1.
                                                                                                                                                                                                                                                          PS50002;
    523 AA;
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                                                                                                                                                                                                                                                          SH3;
    58778
                                                                                                                                                                                                               SH3
                                                                                                                                                                                                             Oncogene; Transferase; Phosphorylation; 5H3 domain; SH2 domain.
    MW;
                                             PROTEIN KINASE.
ATP (BY SIMILARITY).
ATP (BY SIMILARITY).
BY SIMILARITY.
                                                                                                                                                                                     domain; SH2 of MYRISTATE.
                                                                                                                                        SH3.
SH2.
                            PHOSPHORYLATION (AUTO-)
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    853245739F6B90ED CRC64;
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Search completed: July 9, Job time: 205 sec
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GSAB_PICPA
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Best Local :
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Best Local
                                                                                                                                                                                                            EMBL; AF309871; AAG30292.1; -
InterPro; IPR001138; Zn_CY6_fungal.
Pfam; PF00172; Zn_clus; 1
SMART; SM00066; GAL4; 1
SMART; SM00066; ZN_CY6_FUNGAL_2; 1.
Transcription regulation; Activator; DNA-binding; Nuclear protein; Zinc; Metal-binding; Autophagy.
DNA_BIND 632 659
DNA_BIND 632 659
SEQUENCE 1862 AA; 208595 MW; ICC87BECIFA21EF8 CRC64;
                                                                                                                                                                                                                                                                                                                                                                             This SWISS-PROT entry is copyright. It is produced through a collaboration between the Swiss Institute of Bioinformatics and the EMBL outstation the European Bioinformatics Institute. There are no restrictions on its use by non-profit institutions as long as its content is in no way modified and this statement is not removed. Usage by and for commercial entitles requires a license agreement (See http://www.isb-sib.ch/announce/or send an email to license@isb-sib.ch).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       SEQUENCE FROM N.A.

MEDLING-2155190; PubMed-11533052;

Stromhaug P.E., Bevan A., Dunn W.A. Jr.;

"GSAll encodes a unique 208-kDa protein required for pexophagy and autophagy in Pichia pastoris.";

J. Biol. Chem. 276:42422-42435(2001).

-1- FUNCTION: REQUIRED FOR GLUCOSE-INDUCED PEROXISOME AUTOPHAGY
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      (PEXOPHAGY).
-!- SUBCELLULAR LOCATION: Nuclear (By similarity).
-!- SIMILARITY: CONTAINS 1 ZN(2)-CYS(6), FUNGAL-TYPE BINUCLEAR
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                O9HFR4;
16-OCT-2001 (Rel. 40, Created)
16-OCT-2001 (Rel. 40, Last sequence update)
01-MAR-2002 (Rel. 41, Last annotation update)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Pichia pastoris (Yeast).

Eukaryota; Fungi; Ascomycota; Saccharomycotina; Saccharomycetes; Saccharomycetales; Saccharomycetaceae; Pichia.

NCBI_TaxID-4922;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       GSAB_PICPA
                                                                               447 AMVSE 451
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                                                                                                                                            Local Similarity nes 5; Conserv
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les 5; Conserv
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                                                                                                                                            Conservative
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                                                                                                                                    100.0%; Score 22; DB 1; 1
100.0%; Pred. No. 2.2e+02;
vative 0; Mismatches 0;
           2002, 12:23:20
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